

***Urb1* Cas9-CKO Strategy**

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Project Overview

Project Name

Urb1

Project type

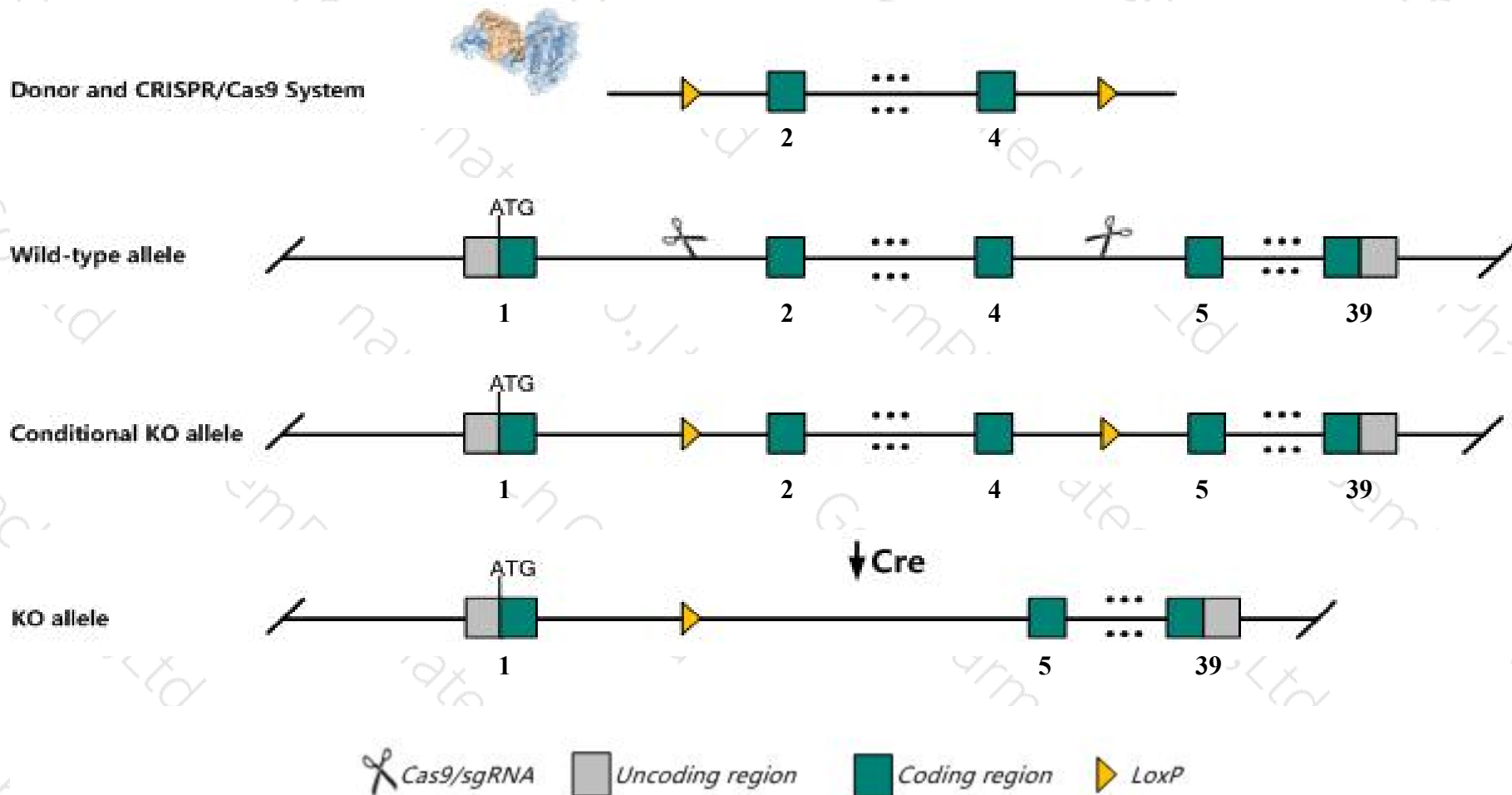
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Urb1* gene. The schematic diagram is as follows:



Technical routes

- The *Urb1* gene has 5 transcripts. According to the structure of *Urb1* gene, exon2-exon4 of *Urb1*-203(ENSMUST00000140920.1) transcript is recommended as the knockout region. The region contains 425bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Urb1* gene. The brief process is as follows: sgRNA was transcribed in vitro, donor vector was constructed. Cas9, sgRNA and Donor were microinjected into the fertilized eggs of C57BL/6JGpt mice. Fertilized eggs were transplanted to obtain positive F0 mice which were confirmed by PCR and sequencing. A stable F1 generation mouse model was obtained by mating positive F0 generation mice with C57BL/6JGpt mice.
- The flox mice was knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

Notice

- The *Urb1* gene is located on the Chr16. If the knockout mice are crossed with other mice strains to obtain double gene positive homozygous mouse offspring, please avoid the two genes on the same chromosome.
- Transcript *Urb1*-202 may not be affected.
- This strategy is designed based on genetic information in existing databases. Due to the complexity of biological processes, all risk of loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)

Urb1 URB1 ribosome biogenesis 1 homolog (S. cerevisiae) [Mus musculus (house mouse)]

Gene ID: 207932, updated on 13-Mar-2020

Summary



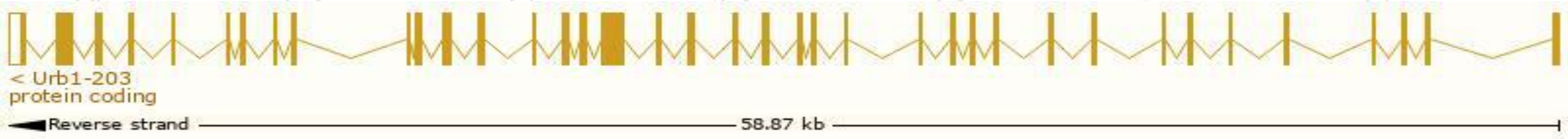
Official Symbol	Urb1 provided by MGI
Official Full Name	URB1 ribosome biogenesis 1 homolog (S. cerevisiae) provided by MGI
Primary source	MGI:MGI:2146468
See related	Ensembl:ENSMUSG00000039929
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	4921511H13Rik, 5730405K23Rik, AA670781, AI645590, mKIAA0539
Expression	Ubiquitous expression in testis adult (RPKM 8.9), CNS E11.5 (RPKM 3.1) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

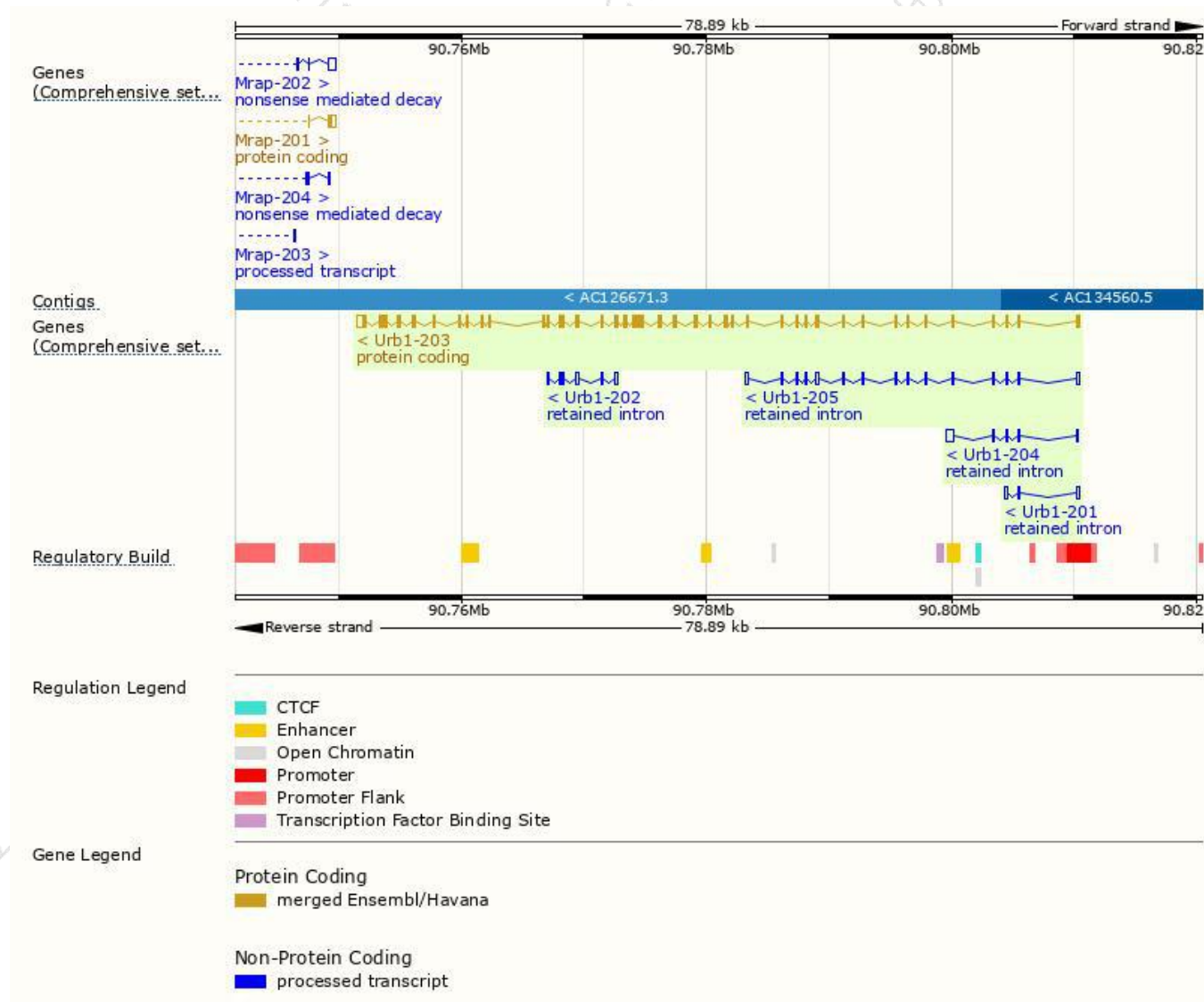
The gene has 5 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Urb1-203	ENSMUST00000140920.1	7344	2277aa	Protein coding	CCDS49905	E9PU96	TSL:5 GENCODE basic APPRIS P1
Urb1-205	ENSMUST00000142955.7	2183	No protein	Retained intron	-	-	TSL:1
Urb1-204	ENSMUST00000140942.7	1218	No protein	Retained intron	-	-	TSL:1
Urb1-202	ENSMUST00000140158.1	809	No protein	Retained intron	-	-	TSL:2
Urb1-201	ENSMUST00000128989.1	661	No protein	Retained intron	-	-	TSL:3

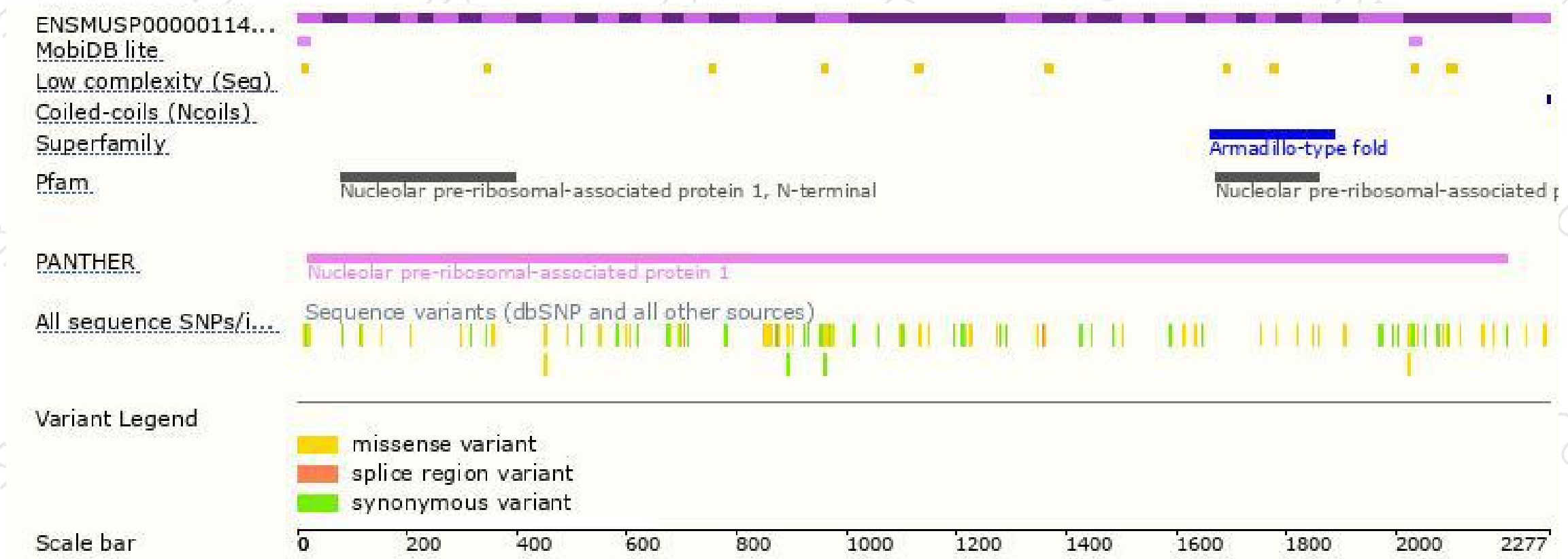
The strategy is based on the design of *Urb1-203* transcript,the transcription is shown below:



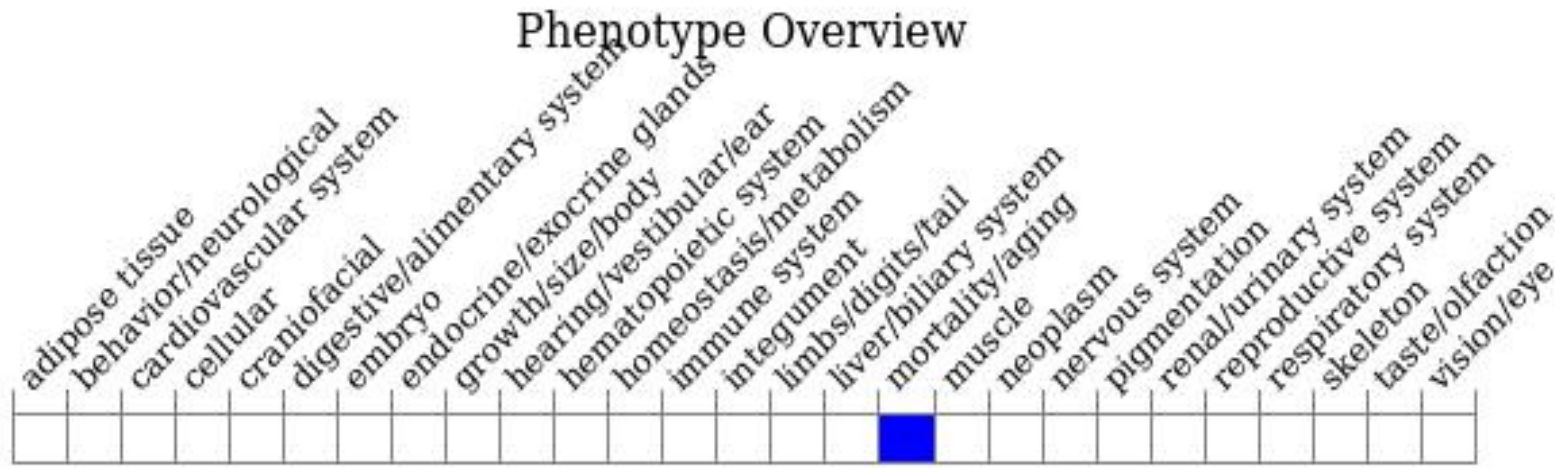
Genomic location distribution



Protein domain



Mouse phenotype description(MGI)



Phenotypes affected by the gene are marked in blue. Data quoted from MGI database(<http://www.informatics.jax.org/>).

If you have any questions, you are welcome to inquire.

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